

FIG. 1

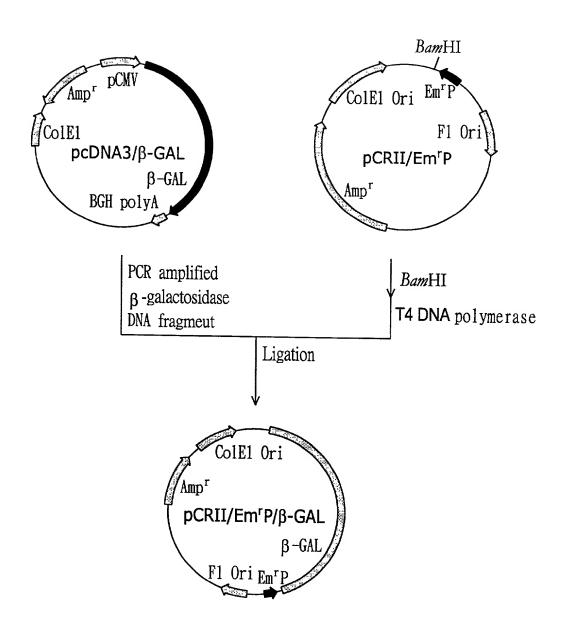
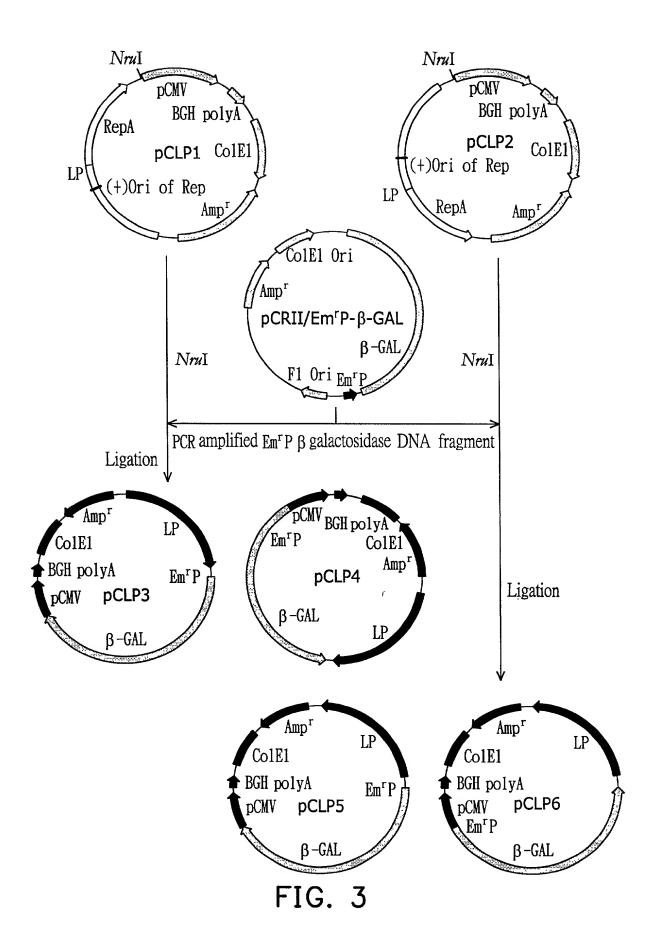


FIG. 2



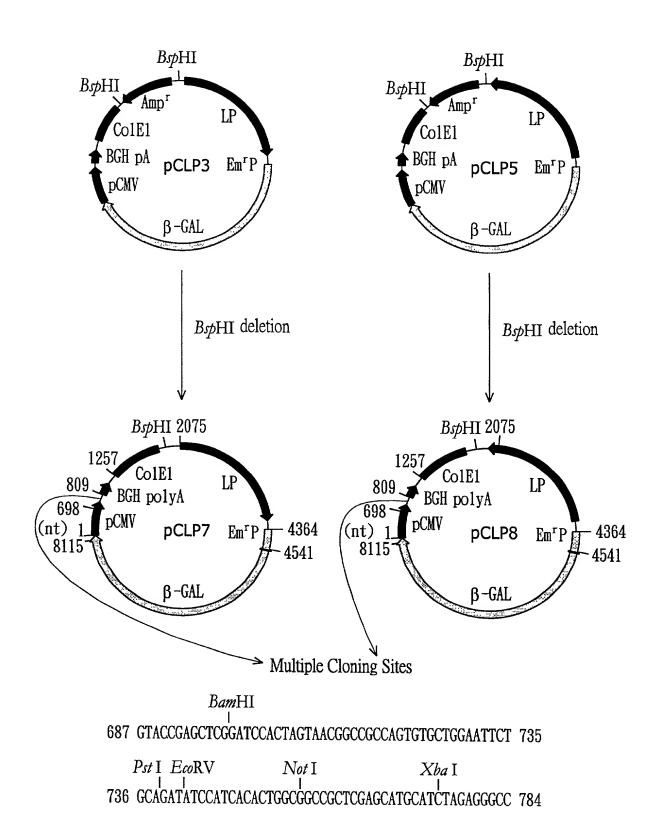


FIG. 4

10	20	30	40	50	60	
GATGTACGGG	CCAGATATAC	GCGTTGACAT	TGATTATTGA	CTAGTTATTA	ATAGTAATCA	
70	80	90	100	110	120	
ATTACGGGGT	CATTAGTTCA	TAGCCCATAT	ATGGAGTTCC	GCGTTACATA	ACTTACGGTA	
130	140	150	160	170	180	
AATGGCCCGC	CTGGCTGACC	GCCCAACGAC	CCCCGCCCAT	TGACGTCAAT	AATGACGTAT	
190	200	210	220	230	240	
GTTCCCATAG	TAACGCCAAT	AGGGACTTTC	CATTGACGTC	AATGGGTGGA	CTATTTACGG	
250	260	270	280	290	300	
TAAACTGCCC	ACTIGGCAGT	ACATCAAGTG	TATCATATGC	CAAGTACGCC	CCCTATTGAC	
310	320	330	340	350	360	
GTCAATGACG	GTAAATGGCC	CGCCTGGCAT	TATGCCCAGT	ACATGACCTT	ATGGGACTTT	
370	380	390	400	410	420	
CCTACTTGGC	AGTACATCTA	CGTATTAGTC	ATCGCTATTA	CCATGGTGAT	GCGGTTTTGG	
430	440	450	460	470	480	
CAGTACATCA	ATGGGCGTGG	ATAGCGGTTT	GACTCACGGG	GATTTCCAAG	TCTCCACCCC	
490	500	510	520	530	540	
ATTGACGTCA	ATGGGAGTTT	GTTTTGGCAC	CAAAATCAAC	GGGACTTTCC	AAAATGTCGT	
550	560	570	580	590	600	<u></u>
AACAACTCCG	CCCCATTGAC	GCAAATGGGC	GGTAGGCGTG	TACGGTGGGA	GGTCTATATA	
610	620	630	640	650	660	FIG. 5
AGCAGAGCTC	TCTGGCTAAC	TAGAGAACCC	ACTGCTTACT	GGCTTATCGA	AATTAATACG	
670	680	690	700	710	720	FIG. 5
ACTCACTATA	GGGAGACCCA	AGCTTGGTAC	CGAGCTCGGA	TCCACTAGTA	ACGGCCGCCA	
730	740	750	760	770	780	rib.
GTGTGCTGGA	ATTCTGCAGA	TATCCATCAC	ACTGGCGGCC	GCTCGAGCAT	GCATCTAGAG	
790 GGCCCTATTC	800 TATAGTGTCA	810 CCTAAATGCT	820 'AGAGCTCGCT	830 GATCAGCCTC	840 GACTGTGCCT	i FIG. !
850	860	870	880	890	900	i
TCTAGTTGCC	AGCCATCTGT	TGTTTGCCCC	TCCCCCGTGC	CTTCCTTGAC	CCTGGAAGGT	
910	920	930	940	950	960	FIG.
GCCACTCCCA	CTGTCCTTTC	CTAATAAAT	GAGGAAATTG	CATCGCATTO	TCTGAGTAGG	
970	980	990	1000	1010	1020	<u> </u>
TGTCATTCTA	TTCTGGGGG	G TGGGGTGGGG	CAGGACAGCA	AGGGGGAGGA	A TTGGGAAGAC	
		1050 TGCGGTGGGC			1080 A AAGAACCAGC	FIG.
TGCATTAATG	AATCGGCCAA	CGCGCGGGGA	A GAGGCGGTTT	GCGTATTGGC	1140 G CGCTCTTCCG	FIG.
CTTCCTCGCT	CACTGACTCC) 1170 CTGCGCTCGG	TCGTTCGGC) 119(F GCGGCGAGCC	1200 G GTATCAGCTC	
			FIG. 5	. A		FIG.

FIG. 5A

1210	1220	1230	1240	1250	1260
ACTCAAAGGC GO	FTAATACGG TT	ATCCACAG A	AATCAGGGGA'	TAACGCAGGA A	AGAACATGT
1270	1280	1290	1300	1310	1320
GAGCAAAAGG CO	CAGCAAAAG GC	CAGGAACC (TAAAAAGGC	CGCGTTGCTG G	CGTTTTTCC
ATAGGCTCCG C	CCCCCTGAC GA	GCATCACA	AAAATCGACG		GGTGGCGAA
1390	1400	1410	1420	1430	1440
ACCCGACAGG A	CTATAAAGA TA	ACCAGGCGT	TTCCCCCTGG	AAGCTCCCTC	FIGCGCTCTC
1450	1460	1470	1480	1490	1500
CTGTTCCGAC C	CTGCCGCTT A	CCGGATACC	TGTCCGCCTT	TCTCCCTTCG (EGAAGCGTGG
1510	1520	1530	1540	1550	1560
CGCTTTCTCA A	ATGCTCACGC TY	GTAGGTATC	TCAGTTCGGT	GTAGGTCGTT	CGCTCCAAGC
1570	1580	1590	1600	1610	1620
TGGGCTGTGT	CACGAACCC C	CCGTTCAGC	CCGACCGCTG	CGCCTTATCC	GGTAACTATC
1630	1640	1650	1660	1670	1680
GTCTTGAGTC (CAACCCGGTA A	GACACGACT	TATCGCCACT	GGCAGCAGCC	ACTGGTAACA
1690	1700	1710	1720	1730	1740
GGATTAGCAG	AGCGAGGTAT G	TAGGCGGTG	CTACAGAGTT	CTTGAAGTGG	TGGCCTAACT
1750	1760	1770	1780	1790	1800
ACGGCTACAC	TAGAAGGACA (TATTIGGTA	TCTGCGCTCT	GCTGAAGCCA	GTTACCTTCG
1810	1820	1830	1840	1850	1860
GAAAAAGAGT	TGGTAGCTCT	IGATCCGCA	AACAAACCA	CGCTGGTAGC	GGTGGTTTTT
1870	1880	1890	1900	1910	1920
TTGTTTGCAA	GCAGCAGATT	ACGCGCAGAA	AAAAAGGATY	TCAAGAAGAT	CCTTTGATCT
1930	1940	1950) 1960	0 1970	1980
TTTCTACGGG	GTCTGACGCT	CAGTGGAACO	AAAACTCAC	G TTAAGGGATT	TIGGTCATGA
1990	2000	2010) 202	0 2030	2040
GCGGATACAT	ATTTGAATGT	ATTTAGAAA	A ATAAACAAA	T AGGGGTTCCG	CGCACATTTC
2050	2060	2070	0 208	0 2090	2100
CCCGAAAAGT	GCCACCTGAC	GTCGACGGA	r CGGGAGATC	A ACGGTAAATC	CGTTGGCATA
2110	2120	213	0 214	0 2150	2160
TCCCTTTTT	GTTGTCAGCT	TGCTGACTT	C TGATACAGO	T TTTAGCATTA	CTCCAATTTA
2170	2180	219	0 220	00 2210) 2220
TTTGGAGTGT	AAGTGCACAT	TATCATGTA	G TGCGCATTA	AT CATGTAGTG	C GCATTATCAT
2230	2240	225	0 226	50 2270	2280
GTAGTGCGCA	TTATCATGTA	GTGCGCATT	A TCATGTAG	NG CGCATTATC	A TGTAGTGCGC
ATTATCATGI		TTATCATGT	'A CATTATCA'	IG TAGTGCGCA	0 2340 T TATCATGTAG
2350) 2360	237	70 23	80 239	0 2400
TGCGCACATT	P ATCATGTAGT	GCGCATTAT	FC ATGTAGTG	CG CATTATCAT	G TAGTGCGCAC

FIG. 5B

		2430 TGTTGTGCTA			
		2490 TGCGTCAGTT			
		2550 TTATTTTTGA			
2590 CTAAGCGATT	2600 TTAGCTAACA	2610 GTTAGCTATC	2620 TAACTGTCTG	2630 TCAACGGTAA	2640 ATCGACTTAG
		2670 AGGCGATATT			
2710 AGACTAGACA	2720 ATACCAAAAG	2730 CCTGACGTCT		2750 GCCCTTGTTT	
		2790 TCCAGCTGGT			
2830 TTTAAAAAAA		2850 GGTTTTTGCT		2870 TTGGGGTTTA	
		2910 CTTGATACTA		2930 CTAGTTTTT	
		2970 TTGTATCCAT			
3010 GGTTATAAAT		3030 AAAGACAAAA		3050 CGTGCAAATT	
		3090 GATTATCATT			3120 AGGGGAAAAG
3130 CCCTATGATO		3150 AAGCTTATTG			3180 CACGTGGATA
					3240 GGCAAAGTTA
3250 GACCTTGGC	3260 G AGAACGGAA	3270 G ATTGAAAATG	3280 TGCGCTATG	3290 C CGAATATTIC	3300 G GCAATCTTAG
					3360 CCTAAGATTG
					3420 F CCATTGTGTA
					0 3480 G GAAGCAGTTG
					0 3540 C GCTCATTCAG
					0 3600 G CTAACTCGCT

FIG. 5C

	3620 GACTAAAAAT			3650 AACGGAAATT A	3660 ACCGTTAATG
	3680 GTCATATAAT			3710 GTTTGTAAAA	
	3740 AAATAATTAT			3770 AAAATTATGG (3780 CAAAAAGCCT
3790 TGAAAGTTGA		3810 GTGGTGCATG		3830 TAAAGCTAAC	
				3890 ATACGAGGTA	
	3920 GGCTGATGAT			3950 TAAAAATTTG	
				4010 GCAAATTAAG	
	4040 TGTTGAGAAT			4070 CGATGAAGAT	
				4130 TTTTAATAAA	
4150 TTATTIGGTA				4190 CCCTATGGTC	
4210 ACAATCTGCT				4250 TCCCTGCTTG	
4270 GTCGCTGAGT				4310 GGCAAGGCTT	
4330 TGCATGAAGA				4370 TTCGTTAGAA	
4390 AGTGTGTTGA				4430 AGGAATTGAA	
				4490 TGGCAGCCAG	
ATTAATGAAC	TTGGACATGO	TTGACGACCC	GGTCTTTGCA	4550 AGCCGAATTC	GACCACACTG
				4610 GACGGCTTTG	
) 4670 CCCCACCTGG	
GCACTCGAT	C GTCCTGGAA	A ATGAACATG	GGTCTTGGT) 4730 C TTGATTCACC	TTGGCCTGGG
475 CACGGTCAA	0 4760 A TTAAACGGG	0 477 A CIGGCITIG	0 4780 I' CAGCTATGT	0 4790 r gaagaggge	4800 GCCAGGTAGA

FIG. 5D

4810 AGCCGGCCAG			4840 CCCGGCGATC		4860 AGCTGGACGA
			4900 AACTTTCACA		
4930 GATCGGCCAC		4950 CCCTGGATGA	4960 TGTATTCAAG	4970 TTAGAAGGGA	
			5020 AGTTGACCAG		
5050 TGACCCGGAA			5080 TCCCCCGCAC		
			5140 CAGTTTAGTG		
5170 GCTGATTGAC			5200 ACCAGTCAAC		
			5260 CGGCAACCTG		
			5320 GGACGGCAGT		
5350 AATTCCAAGC	5360 AAAAATCCGC	5370 TCGCTTCTTA	5380 TGTCAGATAC	5390 TTTGACCTGG	5400 ATGAAGCTTT
			5440 CGGGGCGCA		
5470 GAACGGCCAC			5500 CTCCTTTACC		
			5560 GGCAGTGGCT		
5590 CTCCTGGCTC) 5620 CATGTCTGGT		
) 5680 CCTTAAGCTT		
) 5740 CAATATTGCC		
) 5800 G TGACTTGGT		
			0 5860 C TGATTTGCC		
			0 5920 A TTTATACCA		
			0 5980 T TGAACTAAA		

FIG. 5E

6010 CGGCCAGCGG	6020 ATCGTCTTCA	6030 AGGGGGCCAA	6040 CCGGCACGAA	6050 TTTGACAGTA	6060 AGTTGGGTCG
				6110 ATGAAGCGAA	
				6170 TACCGGCTCT	
6190 CGGCCTTTAC	6200 GTCATTGATG	6210 AAGCTAACCT	6220 GGAAAGCCAC	6230 GGCACCTGGG	6240 AAAAAGTGGG
				6290 CAGCATTGGC	
6310 CTTATCCCGG				6350 CATGCTTCAA	
				6410 CAAATGGCTG	
6430 GAAGGCTGAT				6470 ACCCACAACC	
6490 CGACGCCACC	6500 CAGATTGAA) 6510 GCCGGATGTA	6520 TGCTCCGGCC	6530 AAGGTAATTG	6540 AAGAATACTT
6550 GACCAATAAA				6590 GCTCACGCCA	
6610 CGTCGGTGAC) 6620 C CIGGCCGCC) 6630 r acacggccci) 6640 GGAAAAATAC	6650 CCCCACTACC	6660 AGGGCGGCTT
667(CATCTGGGA() 668 C TGGATTGAC	0 6690 C AAGGACTGGZ) 6700 A AAAAGACGGG	6710 CACCIGCIT	6720 ATGGGGGCGA
673(CTTCGATGA() 6770 GGCCTGGTCT	
679 GACTGAATC	0 680 G CCGAAACTG	0 681 G CTAATGTCA	0 6820 A GGCCCTTTAC	O 6830 C GCCAACCTTA	6840 AGTTAGAAGT
685 AAAAGATGG	0 686 G CAGCTCTIC	0 687 C TCAAAAACG	0 6880 A CAATTTATT	0 6890 T ACCAACAGCT	6900 CATCTTACTA
691 CTTCTTGAC	0 692 T AGTCTTTTC	0 693 G TCGATGGCA	0 694 A GTTGACCTA	0 6950 C CAGAGCCGGO	6960 CTCTGACCTT
697 TGGCCTGGA	0 698 G CCTGGCGAA	0 699 T CCGGGACCT	0 700 T TGCCCTGCC	0 7010 T TGGCCGGAAG) 7020 G TCGCTGATGA
					0 7080 C CTTGGGCGGA
					0 7140 G AATTTAAGCC
715 GGAAGGGC	50 710 FG CCAGATTT	50 717 AG TIGATICCO	70 718 SA CTACAACCI	0 719 A GGCCTGAAA	0 7200 G GAAATAACTI

FIG. 5F

7210 CCAAATICIC	7220 TTCTCCAAGG	7230 TCAAGGGCTG	7240 GCCGGTTTCC	7250 CTCAAGTATG	7260 CCGGTAGGGA
7270 ATACTTGAAG	7280 CGGCTGCCGG	7290 AATTTACCTT	7300 CTGGCGGGCC	7310 CTGACGGACA	7320 ACGACCGGGG
7330 AGCTGGTTAC	7340 GGCTATGATC	7350 TGGCCCGGTG	7360 GGAAAATGCC	7370 GGCAAGTATG	7380 CCCGCTTGAA
7390 AGACATCAGC	7400 TGCGAGGTCA	7410 AGGAAGACTC	7420 CGTTTTGGTC	7430 AAGACTGCCT	7440 TTACGTTGCC
				7490 GATGGACGGG	
7510 TGTAACAGCT				7550 TTGCCAGCCT	
7570 CCTGGCCCTG	7580 CCAAAAGAAC	7590 TGACCGATTA	7600 CCGCTACTAT	7610 GGTCTGGGAC	7620 CTAATGAGAG
				7670 CAGGGAGCGG	
7690 CTTTAGCCCA	7700 TATOGTCCGC	7710 AGGAAACGG	7720 CAACCGGAGG	7730 AAGGTTCGCT	7740 GGTACCAGCT
7750 CTTTGATGAA	7760 AAGGGCGGC1	777(TGGAATTTA	7780 C GGCCAATGG	7790 G GCAGACTIGA	7800 ACTTGTCTGC
7810				0 7850 C GCTTTTGAAC	7860 TGACTAACAA
7870 TTACACTTGG					7920 ACTCCTGGG
7930 GCAGAAGGIY					7980 TTCGCCTGGT
					8040 8040 E
					0 8100 C AGCCGGCGTG
811 TCGCTAGCG	0 812 A CGGTC	0 813	814	10 815	0 8160

10	20	30	40	50	60	
GATGTACGGG	CCAGATATAC	CCCTTGACAT	TGATTATTGA	CTAGTTATTA	ATAGTAATCA	
	80 CATTAGTTCA					
130	140	150	160	170	180	
AATGGCCCGC	CTGGCTGACC	GCCCAACGAC	CCCCGCCCAT	TGACGTCAAT	AATGACGTAT	
190	200	210	220	230	240	
GTTCCCATAG	TAACGCCAAT	AGGGACTTTC	CATTGACGTC	AATGGGTGGA	CTATTTACGG	
250	260	270	280	290	300	
TAAACTGCCC	ACTTGGCAGT	ACATCAAGTG	TATCATATGC	CAAGTACGCC	CCCTATTGAC	
310	320	330	340	350	360	
GTCAATGACG	GTAAATGGCC	CGCCTGGCAT	TATGCCCAGT	ACATGACCTT	ATGGGACTTT	
370	380	390	400	410	420	
CCTACTTGGC	AGTACATCTA	CGTATTAGTC	ATCGCTATTA	CCATGGTGAT	GCGGTTTIGG	
430	440	450	460	470	480	
CAGTACATCA	ATGGGCGTGG	ATAGCGGTTT	GACTCACGGG	GATTTCCAAG	TCTCCACCCC	
490	500	510	520	530	540	
ATTGACGTCA	ATGGGAGTTT	GTTTTGGCAC	CAAAATCAAC	GGGACTTTCC	AAAATGTCGT	
550	560	570	580	590	600	L
AACAACTCCG	CCCCATTGAC	GCAAATGGGC	GGTAGGCGTG	TACGGTGGGA	GGTCTATATA	
610 AGCAGAGCTC	TCTGGCTAAC	TAGAGAACCC	ACTGCTTACT	GGCTTATCGA	660 AATTAATACG	
670	680	690	700	710	720	FIG. 6B
ACTCACTATA	GGGAGACCCA	AGCTTGGTAC	CGAGCTCGGA	TCCACTAGTA	ACGGCCGCCA	
GTGTGCTGGA	ATTCTGCAGA	TATCCATCAC	: ACTGGCGGC	GCTCGAGCAT		FIG. 00
GGCCCTATIC	TATAGTGTCA	CCTAAATGCT	AGAGCTCGC	GATCAGCCIV	840 GACTGTGCCT	1
850) 860	870) 886) 89(900	
TCTAGTTGC0	C AGCCATCTGT	GTTTGCCCC	C TCCCCCGTG	CTTCCTTGA(CCTGGAAGGT	
910) 920	930	94	950	960	FIG. 6D
GCCACTCCC	A CTGTCCTTTC	CTAATAAAA	F GAGGAAATTV	G CATCGCATIV	G TCTGAGTAGG	
970 TGTCATTCTX				0 1010 A AGGGGGAGG	0 1020 A TTGGGAAGAC	<u> </u>
	0 1040 C ATGCTGGGG			0 107 T CTGAGGCGG	0 1080 A AAGAACCAGC	FIG. 6E
	G AATCGGCCA	A CGCGCGGGG	A GAGGCGGTT		G CGCTCTTCCG	FIG. 6F
115 CTTCCTCGC	0 116 T CACTGACTO	0 117 G CTGCGCTCG	0 118 G TCGTTCGGC		0 1200 G GTATCAGCTC	1
						FIG. 6G
			FIG. 6	δA		L

1210 ACTCAAAGGC	1220 GGTAATACGG	1230 TTATCCACAG	1240 AATCAGGGGA	1250 TAACGCAGGA	1260 AAGAACATGT
1270 GAGCAAAAGG	1280	1290	1300 GTAAAAAGGC	1310 CGCGTTGCTG (1320
1330	1340	1350	1360	1370	1380
ATAGGCTCCG	CCCCCTGAC	GAGCATCACA	AAAATCGACG	CTCAAGTCAG	AGGTGGCGAA
1390 ACCCGACAGG			1420 TTCCCCCTGG		
			1480 TGTCCGCCTT		
1510 CGCTTTCTCA			1540 TCAGTTCGGT		
			1600 CCGACCGCTG		
			1660 TATCGCCACT		
1690 GGATTAGCAG	1700 AGCGAGGTAT	1710 GTAGGCGGTG	1720 CTACAGAGTT	1730 CTTGAAGTGG	1740 TGGCCTAACT
	1760 TAGAAGGACA		1780 TCTGCGCTCT		
1810 GAAAAAGAGI	1820 TGGTAGCTCT	1830 TGATCCGGCA	1840 AACAAACCAC	1850 CGCTGGTAGC	1860 GGTGGTTTT
1870 TTGTTTGCAA			1900 AAAAAGGATC		
1930) 1960 S AAAACTCACG		
1990 GCGGATACA) 2020 A ATAAACAAAT		
2050 CCCGAAAAG) 2080 I CGGGAGATCA		
			D 2140 C CATTTGCGA		
			0 2200 A ACATGAACTA		
			0 2260 T AATAATCCAC		
			0 2320 C ACCACCAAA		
235 ATATAATCA			0 2380 C GCCGTTTCT		

FIG. 6C

AGCCCCTCTA AGTCGATTTA CCGTTGACAG ACAGTTAGAT AGCTAACTGT TAGCTAAAAT CGCTTAGAAC GCAAATAAGA GCCTTTAAAA TTAACGTTCA AAAATAAAAA AGTTCGAAGG AGCTAGCGAC TGAACTTATT TATTTTTGAA TGTTCCAAAC TGACGCAAGT CAGTTACGTT TGTGTAAGTG CGCACTACAT GATAATGCGC ACTACATGAT AATGCGCACT ACATGATAAT GTGCGCACTA CATGATAATG CGCACTACAT GATAATGTAC ATGATAATGT GCGCACTACA TGATAATGCG CACTACATGA TAATGCGCAC TACATGATAA TGCGCACTAC ATGATAATGC GCACTACATG ATAATGCGCA CTACATGATA ATGCGCACTA CATGATAATG TGCACTTACA CTCCAAATAA ATTGGAGTAA TGCTAAAACC TGTATCAGAA GTCAGCAAGC TGACAACAAA AAAGGATAT GCCAACGAT TTACCGTTGA TCTCCCGATC CCCTATGGTC GACTCTCAGT ACAATCTGCT CTGATGCCGC ATAGTTAAGC CAGTATCTGC TCCCTGCTTG TGTGTTGGAG GTCGCTGAGT AGTGCGCGAG CAAAATTTAA GCTACAACAA GGCAAGGCTT GACCGACAAT TGCATGAAGA ATCTGCTTAG GGTTAGGCGT TTTGCGCTGC TTCGTTAGAA GCAAACTAAG AGTGTGTGA GTAGTGCAGT ATCTTAAAAT TTTGTATAAT AGGAATTGAA GTTAAATTAG ATGCTAAAAA TTTGTAATTA AGAAGGAGTG ATTACATGAT TGGCAGCCAG TCTCCGGGCA ATTAATGAAC TIGGACATGG TIGACGACCC GGTCTTIGCA AGCCGAATTC GACCACACTG GCGCCGTTA CTAGGGTATC GATCCGATAA AAAGTTAGGC GACGCCTTTG CCCTGGTGCC AGCAGACGGT AAGGTCTACG CGCCATTTGC CGGTACTGTC CGCCAGCTGG CCAAGACCCG GCACTOGATO GTCCTGGAAA ATGAACATGG GGTCTTGGTC TTGATTCACC TTGGCCTGGG CACGGTCAAA TTAAACGGGA CTGGCTTTGT CAGCTATGTT GAAGAGGGCA GCCAGGTAGA

FIG. 6D

4810 AGCCGGCCAG	4820 CAGATCCTGG				
	4880 GTGACCGTCA				
4930 GATCGGCCAC	4940 AGCGTCCAAG	4950 CCCTGGATGA	4960 TGTATTCAAG	4970 TTAGAAGGGA	4980 AGAATTAGAA
	5000 AAGTTAGTAA				
	5060 GTTTACGAAG				
	5120 GAACTGGAGG				
5170 GCTGATTGAC	5180 TACGCTGAAA	5190 ACGGCCAGGG	5200 ACCAGTCAAC	5210 TTCTATGCAG	5220 AAGACTTTGA
	5240 TTTAAGTCAG			5270 GAACTGCAAG	
5290 GCCCCAGTAT	5300 GTCAACGTCC			5330 GAGGAGATTT	
5350 AATTCCAAGC	5360 AAAAATCCGC			5390 TTTGACCTGG	
5410 CTGGGACAAG	5420 GAAGTCAGCT			5450 ACAGCCATCT	
5470 GAACGGCCAC	5480 TTCGTCGGCT			5510 CCAAGCGAGT	
5530 CAAGTTCCTC	5540 AAGAAAGAAA	5550 ATAACCGCCT	5560 GGCAGTGGCT	5570 CTCTACAAGI	5580 ATTCTTCCGC
5590 CTCCTGGCTG				5630 TTGTTCAGAT	5640 CAGTGACTCT
5650 TCAGGCCAAC) 5660 CCGCCTCTGC	5670 ACTIGGAGGA	5680 CCTTAAGCT) 5690 ACGGCCAGCI	5700 TGACCGATAA
		5730 AAGTCGAAGC			5760 CAAATGCCAG
5770 CTTTAAGCIY) 5780 GAAGTGCGGC	5790 ATAGTGAAG) 5800 G TGACTTGGT) 5810 F GCTGAAAAG	5820 TGGGCCCAAT
) 5880 r ggagcgcgga
	0 5900 C CTTTACCAGO				0 5940 C TCTTAGAGGT
595 TAGCCGGCA	0 5960 G GAAGTGGGT) 5970 T TCCGCAACT	0 598 I TGAACTAAA	0 599 A GACGGGATT	0 6000 A TGTACCTTAA

FIG. 6E

6010 CGGCCAGCGG	6020 ATCGTCTTCA	6030 AGGGGGCCAA	6040 CCGGCACGAA	6050 TTTGACAGTA	6060 AGTTGGGTCG
	6080 GAAGAGGATA				
	6140 TGCTCTCACT				
	6200 GTCATTGATG				
	6260 GATCCTAGCT				
	6320 GTGAAGAACA				
	6380 AATGAGTCTT				
GAAGGCTGAT	6440 CCGACCCGGG	TTCAGCACTA	TGAAGGGGTG	ACCCACAACC	GGAAGTTTGA
	6500 CAGATTGAAA				
	6560 CCAGCCAAGC	CATTTATCTC	AGTTGAATAC	GCTCACGCCA	TGGGCAACTC
6610 CGTCGGTGAC	6620 CTGGCCGCCT	6630 ACACGGCCCT	6640 GGAAAAATAC	6650 CCCCACTACC	AGGGCGCTT
6670 CATCTGGGAC	6680 TGGATTGACC	6690 AAGGACTGGA			
6730 CTTCGATGAC	6740 CGGCCAACCG	6750 ACTATGAATT	6760 CTGCGGGAAC	6770 GGCCTGGTCT	6780 TTGCTGACCG
	CCGAAACTGG		GGCCCTTTAC	C GCCAACCTTA	AGTTAGAAGT
6850 AAAAGATGG	6860 CAGCTCTTCC	6870 TCAAAAACGA	6880 CAATTTATT) 6890 F ACCAACAGC	6900 CATCTTACTA
6910 CTTCTTGAC) 6920 r agrettitige	6930 TCGATGGCAA	6940 GTTGACCTAC	O 6950 C CAGAGCCGGC	6960 C CTCTGACCTT
					7020 TCGCTGATGA
AAAAGGGGA	G GTCGTCTAC	C GGGTAACGG(C CCACTTAAA	A GAAGACTIG	7080 CCTTGGGCGGA
TGAGGGCTT	C ACTGTGGCTY	G AAGCAGAAGA	A AGTAGCTCA	A AAGCTGCCG	0 7140 G AATTTAAGCC
					0 7200 G GAAATAACTT

FIG. 6F

7210	7220	7230	7240	7250	7260
				CTCAAGTATG	
7270	7280	7290	7300	7310	7320
ATACTIGAAG	CGGCTGCCGG	AATTTACCTT	CTGGCGGGCC	CTGACGGACA	ACGACCGGGG
7330	7340	7350	7360	7370	7380
AGCTGGTTAC	GGCTATGATC	TGGCCCGGTG	GGAAAATGCC	GGCAAGTATG	CCCGCTTGAA
7390	7400	7410	7420	7430	7440
AGACATCAGC	TGCGAGGTCA	AGGAAGACTC	CGTTTTGGTC	AAGACTGCCT	TTACGTTGCC
7450	7460	7470	7480	74 90	7500
				GATGGACGGG	
				7550	
TGTAACAGCT	GACTICCCAG	GCGCGGAAGA	AGCTGGTCTC	TIGCCAGCCT	TTGGCTTGAA
7570	7580	7590	7600	7610	7620
CCTGGCCCTG	CCAAAAGAAC	TGACCGATTA	CCGCTACTAT	GGTCTGGGAC	CTAATGAGAG
7630	7640	7650	7660	7670	7680
				CAGGGAGCGG	
7690	7700	7710	7720	7730	7740
		AGGAAACGGG	CAACCGGAGC	AAGGTTCGCT	GGTACCAGCT
7750	7760	7770	7790	7790	7800
CTTTGATGAA	AAGGGCGGCT	TGGAATTTAC	GGCCAATGGG	GCAGACTIGA	ACTIGICIGC
				7850	
7810					
TTIGCCATAT				GCTTTTGAAC	
7870				7910	
TTACACTTGG	GTTAGAGCCT	' TAAGCGCCCA	A GATGGGGGTC	: GGCGGGGATG	ACTCCTGGGG
7930	7940	7950	7960	7970	7980
GCAGAAGGTC	CACCCGGAAT	TCTGCCTGG	TGCTCAAAAA	A GCCCGCCAGC	TICGCCIGGT
7990					8040 RAATTTTAGA
GATTCAGCCC				_	
8050	8060	8070	808	8090	8100
AAAAGCAAA	G CGAGTGAGG!	A AGATGGCAA	C GATCAGAGAI	A GIGCCAAGG	C AGCCGGCGTG
8110	8120) 813	0 814	0 8150	8160
TCGCTAGCG	A CGGTC				

FIG. 6G

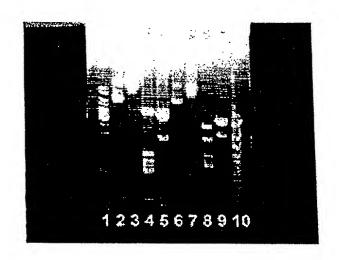


FIG. 7A

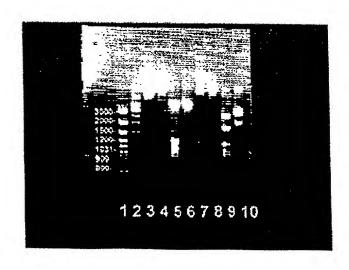


FIG. 7B